

FIG. 1

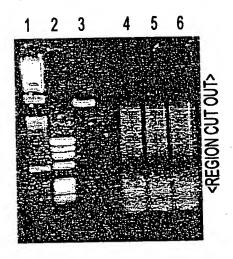


FIG. 2

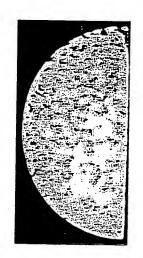
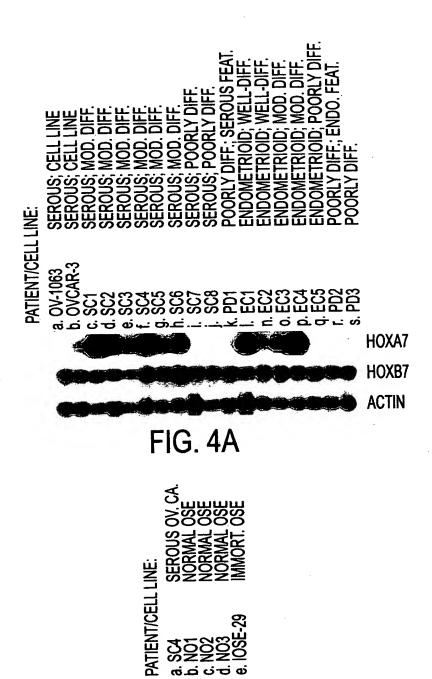


FIG. 3A

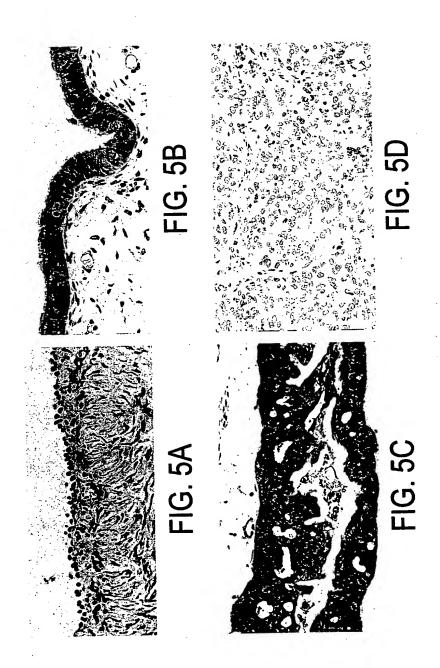


FIG. 3B



HOXA7
HOXB7
ACTIN

FIG. 4B



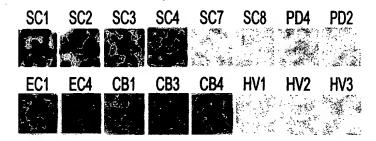
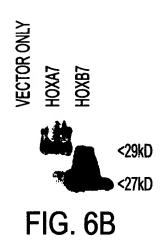


FIG. 6A



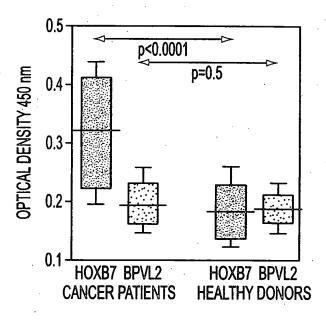
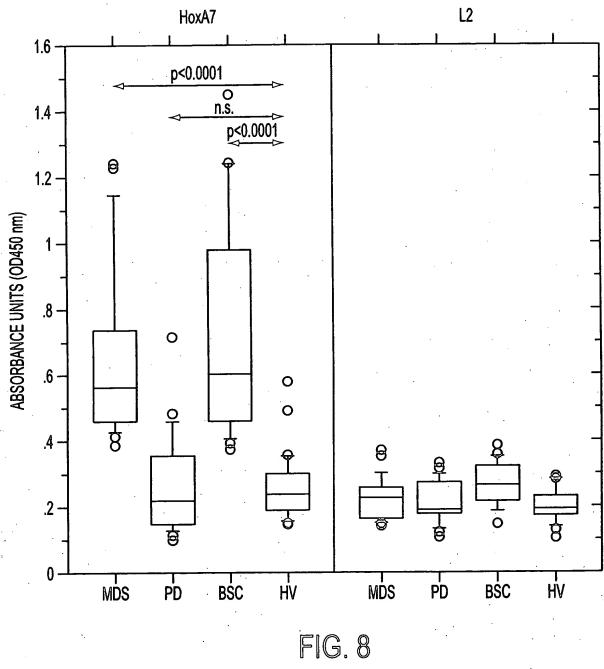


FIG. 7



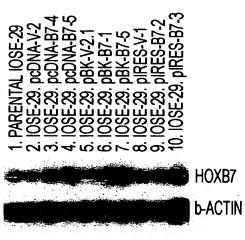


FIG. 9A



FIG. 9B

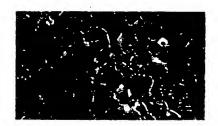
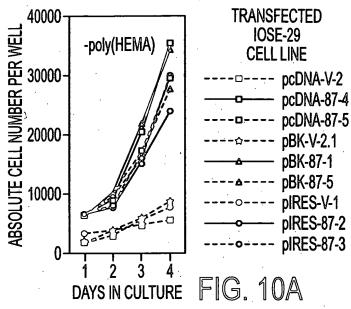
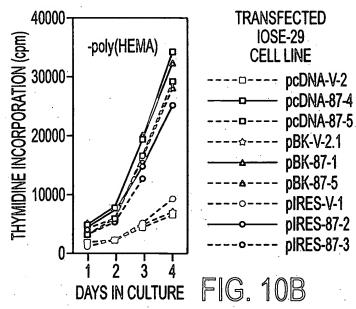
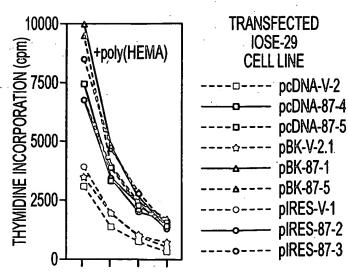


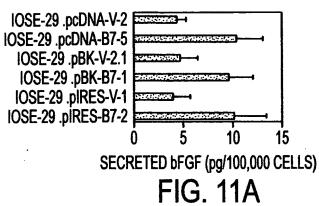
FIG. 9C

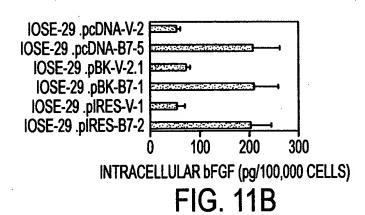












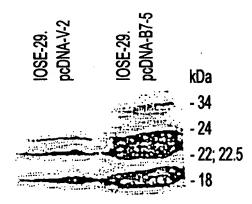


FIG. 11C

atg agt tca ttg tat tat gcg aat gct tta ttt tct aaa tat cca gcc Met Ser Ser Leu Tyr Tyr Ala Asn Ala Leu Phe Ser Lys Tyr Pro Ala tca agt tcg gtt ttc gct acc gga gcc ttc cca gaa caa act tct tgt Ser Ser Ser Val Phe Ala Thr Gly Ala Phe Pro Glu Gln Thr Ser Cys gcg ttt gct tcc aac ccc cag cgc ccg ggc tat gga gcg ggt tcg ggc 144 Ala Phe Ala Ser Asn Pro Gln Arg Pro Gly Tyr Gly Ala Gly Ser Gly gct tcc ttc gcc gcc tcg atg cag ggc ttg tac ccc ggc ggg ggg ggc 192 Ala Ser Phe Ala Ala Ser Met Gln Gly Leu Tyr Pro Gly Gly Gly atg gcg ggc cag agc gcg gcc ggc gtc tac gcg gcc ggc tat ggg ctc 240 Met Ala Gly Gln Ser Ala Ala Gly Val Tyr Ala Ala Gly Tyr Gly Leu 65 70 gag ccg agt tcc ttc aac atg cac tgc gcg ccc ttt gag cag aac ctc 288 Glu Pro Ser Ser Phe Asn Met His Cys Ala Pro Phe Glu Gln Asn Leu 95 tee ggg gtg tgt eee gge gae tee gee aag geg geg gge gee aag gag 336 Ser Gly Val Cys Pro Gly Asp Ser Ala Lys Ala Ala Gly Ala Lys Glu 105 cag agg gac tcg gac ttg gcg gcc gag agt aac ttc cgg atc tac ccc 384 Gln Arg Asp Ser Asp Leu Ala Ala Glu Ser Asn Phe Arg Ile Tyr Pro tgg atg cga agc tca gga act gac cgc aaa cga ggc cgc cag acc tac 432 Trp Met Arg Ser Ser Gly Thr Asp Arg Lys Arg Gly Arg Gln Thr Tyr 130 135 acc cgc tac cag acc ctg gag ctg gag aaa gaa ttt cac tac aat cgc 480 Thr Arg Tyr Gln Thr Leu Glu Leu Glu Lys Glu Phe His Tyr Asn Arg 145 150 155 160 tac ctg acg cgg cgg cgc atc gag atc gcg cac acg ctc tgc ctc 528 Tyr Leu Thr Arg Arg Arg Ile Glu Ile Ala His Thr Leu Cys Leu 170 175 acg gaa aga cag atc aag att tgg ttt cag aac cgg cgc atg aag tgg 576 Thr Glu Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp 180 185 624 Lys Lys Glu Asn Lys Thr Ala Gly Pro Gly Thr Thr Gly Gln Asp Arg 200 gct gaa gca gag gaa gag gaa gag tga gggatggaga aagggcagag 674 Ala Glu Ala Glu Glu Glu Glu Glu 210 215 gaagagacat gagaaaggga gaggaagaga agcccagctc tgggaactga atcaggaaac 734

DOT NOT ENTER GRAJOS